

Boo

Go Clear

## Details

**Send to**

## Links

dbEST Id: 1075499  
EST name: zw78b02.s1  
GenBank Acc: AA432248  
GenBank gi: 2114636

Clone Id: IMAGE:782283 (3')  
Source: IMAGE Consortium, LLNL  
DNA type: cDNA

Sequencing: -41m13 fwd. ET from Amersham  
PolyA Tail: Unknown

TTTTTTTTTTTTTTTTCATTTTGAAAATGCTTTAATAAGTGTTGACAACACTGTTTTGCAA  
AATGTAAAGATACTATACAAATTCTTAATACAAAAAGAATAAATTAAAAGCAGATTTCTT  
TTTTTAATTCTGCAACTTTGTCTACAACGTACATCTTTTTTCATTGATTACAGTTGAACAG  
AATCCAGTAAAAATCATTTTACATGCTCTACAGTCAGTTTCAGGAGCAACCTAATCTTTTT  
TCCCCATTATTAAACTAGAGTCCATTTTACACAACCTTGTAATAAACTATTGACATTAAT  
GTATATGTAAAACTTTACACCTAGTTAACTAAGCAGTAACTGGTCATCTGATAGCACCTG  
GATGGGGTTTGCTATATTTAGAACTAACTAATACTGAATGAAAACAAATTGGAATTTTA  
A

Entry Created: May 22 1997  
Last Updated: May 22 1997

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further information.

```

Lib Name:      Soares_testis_NHT
Organism:      Homo sapiens
Sex:           male
Lab host:      DH10B
Vector:        pT7T3D-Pac (Pharmacia) with a modified polylinker
R. Site 1:     Not I
R. Site 2:     Eco RI
Description:    1st strand cDNA was prepared from mRNA obtained from
                Clontech Laboratories, Inc., and primed with a Not I -
                oligo(dT) primer [5'
                TGTTACCAATCTGAAGTGGGAGCGGCCGCCCAATTTTTTTTTTTTTTTTTT 3'].
                Double-stranded cDNA was ligated to Eco RI adaptors
                (Pharmacia), digested with Not I and cloned into the Not I

```

and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization to Cot5, and was constructed by Bento Soares and M. Fatima Bonaldo.

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

Title: WashU-Merck EST Project 1997  
Authors: Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost  
,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin  
,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising  
,B., White,Y., Wylie,T., Waterston,R., Wilson,R.  
Year: 1997  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



CGCTCAGGATACGACTTCGCTCCTAGAGATCGGATCCCCGGGATATTATTATAGCTCGATCGATCT  
TTCTCTATATCCGCGATGGGATATATACACACACATGCGCGATAGCATGACTGATCT  
CCCCATCT  
CACAGACATACGCT

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

OMIM

Boo

Search  for   

Limits

Preview/Index

History

Clipboard

Details

  Show:   ☐ 1: H18957. ym46g08.s1 Soares...[gi:885197]

Links

## IDENTIFIERS

dbEST Id: 278895  
EST name: ym46g08.s1  
GenBank Acc: H18957  
GenBank gi: 885197  
GDB Id: 424323

## CLONE INFO

Clone Id: IMAGE:51513 (3')  
DNA type: cDNA

## PRIMERS

Sequencing: Promega. -21m13  
PolyA Tail: Unknown

## SEQUENCE

TTTTTTTTTGTTTTAAACACTTTATTTATAAAAAAGTACATTTTTAATCCTCAGTACATT  
TTCAACCCATCATTTTTTTTTTAATACAAGTAAAAGGGGGTGATGCAAACACCCCCCAGGT  
CAGAACCAGGAGGATCTGCTGGGCTGTCCCTGGACCAAAGGCGGAAAGGGCGACAAGACG  
CCGAAGCAAGGTAGCGCATCACGCTGGGAGGGGAGGGTGGCAGCTTCTCCTGGGATTCTT  
TTCATTTATACAAAAAGGAAAACCAATTTTTTCGACCAAGAATCCCATTCTCACAGCA  
GGGGTCAGAAGAGCAGCAGCACCGAGT

Quality: High quality sequence stops at base: 305

Entry Created: Jun 29 1995

Last Updated: Jun 29 1995

## COMMENTS

High quality sequence stops: 305  
Source: IMAGE Consortium, LLNL  
This clone is available royalty-free through LLNL ; contact  
the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further  
information.

## LIBRARY

Lib Name: Soares infant brain 1NIB  
Organism: Homo sapiens  
Sex: female  
Organ: whole brain  
Develop. stage: 73 days post natal  
Lab host: DH10B (ampicillin resistant)  
Vector: Lafmid BA  
R. Site 1: Not I  
R. Site 2: Hind III  
Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer  
[5' AACTGGAAGAATTCGCGCCGAGGAATTTTTTTTTTTTTTTTTT 3'];  
double-stranded cDNA was ligated to Hind III adaptors

(Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafmid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

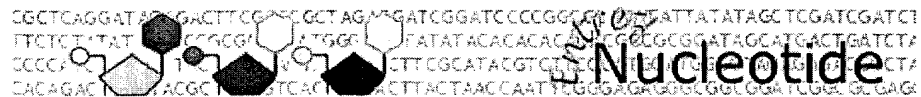
Title: The WashU-Merck EST Project  
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,  
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,  
Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,  
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,  
Wilson,R.  
Year: 1995  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



Boo

Go Clear

## Details

**Send to**

## Links

**dbEST Id:** 4472956  
EST name: EST379537  
GenBank Acc: AW967462  
GenBank qi: 8157299

Plate: 241  
DNA type: cDNA

Sequencing: Forward  
PolyA Tail: Unknown

AACATTAAGATTTTATTACAAACCATGCATTATATATTTCTTTACACTTAAGGAATAGAT  
ATGAAACAATCTTGGAGTAAAAAATTAGAAGGCAACTTGCTTCAAGTTTGTACCAAGTCAA  
TCAAGCAGAAACCTGAAGAACCTTGTTTTAAGATGAGAGTCATTTATACTTGGCAGGCAT  
TTTCTTCCAATGAAAAAATAAAGTCAATGTGCCATTATCTTGACACTTATAAAAAATGTTT  
ATAAAAAGCATTTAGGCCATTGATTCTCACAGTTGGCTGAATATTGGAATCACCTAGATT  
AAAAAAAATACTAATCCCTATACAACATCCCCAAAATTAGATTAAATTAGTGTAAGTTA  
GGCCCTGGGCATATAGGCTGTTTTAAAAATTCTCGGGTGAGTCTAATGTGTACCAGTTTG  
AAAACCACTAGCTTAGAGACCTGTTCCATCATCATAGAAGTCACTCTTTAGCTGAATGAT  
TCTATATGCCCTGATTTTCAGTGGGTAGTGTTGTTATTTTAAACATCTGATAGGTGTATCT  
CCCTTGATTAAAAAATAAATACCTTATGGAAGAGATTATATGGTTTTATTTA

Entry Created: Jun 1 2000  
Last Updated: Jun 1 2000

Lib Name: MAGE resequences, MAGJ  
Organism: Homo sapiens  
Vector: pBluescriptSKm

Name: John Quackenbush  
Institution: The Institute for Genomic Research  
Address: 9712 Medical Center Dr., Rockville, MD 20850, USA  
Tel: 301 838 3528  
Fax: 301 838 0208  
E-mail: johnq@tigr.org

Title: Assessment of gene expression patterns in a model of colon tumor metastasis using a 19,200 element cDNA microarray

Authors: Hegde, P., Qi, R., Abernathy, K., Dharap, S., Gaspard, R., Gay, C., Holt, I.E., Saeed, A.I., Sharov, V., Lee, N.H., Yeatman, T.J.,

Year: Quackenbush, J.  
2000  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



CGCTCAGGATAGGACTTCGCTAGGATCGGATCCCGGCTTATTATATAGCTCGATCGATCT  
TTCTCTATATCCGCGATGGGATATACACACACATCGGCGGATAGCATGACTGATCT  
CCCCATCT  
CACAGACTACGCT

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

OMIM

Boo

Search  for   

Limits

Preview/Index

History

Clipboard

Details

  Show:   ☐ 1: N71027. za35d03.s1 Soares...[gi:1227607]

Links

**IDENTIFIERS**

dbEST Id: 477121  
EST name: za35d03.s1  
GenBank Acc: N71027  
GenBank gi: 1227607  
GDB Id: 1239454

**CLONE INFO**

Clone Id: IMAGE:294533 (3')  
Source: IMAGE Consortium, LLNL  
DNA type: cDNA

**PRIMERS**

Sequencing: m13 -40 forward  
PolyA Tail: Unknown

**SEQUENCE**

AACCTAGACAAAAAGAAGAACCAAGTATAATAAAAAAATCAAGAAACATGACAATAATG  
GAGACTAAATCCCTCGATGGAAAGAATTATAATCATAAATGTAAGCCTTAAAGAGTTAAC  
TTTAGATGTAGACAAGTCAAAGTGAAGCACTAAAAACATGTTTCGACCTTATATAACACAT  
TCCTTAGGGGGAAAAAAGATTTTTTAATATTGGAAAAAATATGGTTAGGCATAGTGGCT  
CATGCCTGTAATCCCAGCACATTCAGAGGCCAAGGCGGGAGGACTGCTTGAGCTTAGGAG  
TTCAAGACCAGCCTGGGCAACATAGTGAGACCTTGTCTCTACTTAAAAACCTAAAAAGAT  
TAGCTGAGTATGGGTGGCATGCACCTATAGTCCCCGGCTACTANGGGAAGCTGAGGGC

Quality: High quality sequence stops at base: 362

Entry Created: Mar 14 1996  
Last Updated: Mar 14 1996

**COMMENTS**

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further information.

PUTATIVE ID Assigned by submitter  
contains Alu repetitive element;

**LIBRARY**

Lib Name: Soares fetal liver spleen 1NFLS  
Organism: Homo sapiens  
Sex: male  
Organ: Liver and Spleen  
Develop. stage: 20 week-post conception fetus  
Lab host: DH10B (ampicillin resistant)  
Vector: pT7T3D (Pharmacia) with a modified polylinker  
R. Site 1: Pac I  
R. Site 2: Eco RI

**Description:** 1st strand cDNA was primed with a Pac I - oligo(dT) primer [5' AACTGGAAGAATTAATTAAAGATCTTTTTTTTTTTTTTTTTTTT 3'], double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

**SUBMITTER**

**Name:** Wilson RK  
**Institution:** Washington University School of Medicine  
**Address:** 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
**Tel:** 314 286 1800  
**Fax:** 314 286 1810  
**E-mail:** [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

**Title:** The WashU-Merck EST Project  
**Authors:** Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P., Wilson, R.  
**Year:** 1995  
**Status:** Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45





Boo

Clear

## Details

File 

## Links

Lib Name: Soares\_NhHMPu\_S1  
Organism: Homo sapiens  
Organ: mixed (see below)  
Tissue type: Pooled human melanocyte, fetal heart, and pregnant uterus  
Lab host: DH10B  
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker  
R. Site 1: Not I

R. Site 2: Eco RI  
Description: Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479.

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

Title: WashU-Merck EST Project 1997  
Authors: Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost ,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin ,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising ,B., White,Y., Wylie,T., Waterston,R., Wilson,R.  
Year: 1997  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



CGCTCAGGATAGGACTTCGTCGCTAGAGATCGGATCCCGGGGATATTATTATAGCTCGATCGATC  
TTCTCTATATTCGCGGATGGGATATATACACACACATCGGCGCGGATAGCATGACTGATCTA  
GCCCATTCTTCGCGGATGGGATATATACACACACATCGGCGCGGATAGCATGACTGATCTA  
CACAGACATACGCTTCGCTAGAGATCGGATCCCGGGGATATTATTATAGCTCGATCGATC

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

OMIM

Bio

Search  for   

Limits

Preview/Index

History

Clipboard

Details

  Show:   ☐ 1: AA620885. af95g06.s1 Soares...[gi:2524824]

Links

## IDENTIFIERS

dbEST Id: 1318692  
EST name: af95g06.s1  
GenBank Acc: AA620885  
GenBank gi: 2524824

## CLONE INFO

Clone Id: IMAGE:1055578 (3')  
Source: IMAGE Consortium, LLNL  
Insert length: 1052  
DNA type: cDNA

## PRIMERS

Sequencing: -40m13 fwd. ET from Amersham  
PolyA Tail: Unknown

## SEQUENCE

TTTGAACATAAAAATTCTTTATTTAACCTAATCCAGCCAGTATTGAGATAGTTTGCTATA  
TTAAAAACAAGACGTTTTAAAAAATTACAGCAAAGTTAGCAAGGCAGTGACTAATTAAGT  
CACTAAGTTTAAATTTTATATTCTTCACAGTCATTTTCATAATCATGTAATGGTAAACAATA  
TTTTTCAGCCACTTTGGAGATAAGTTAACTTTTGAAAAGAATAGAATTCTAGTAGTCGTCA  
TTGAATTTTATAAAAGAGGTTTAAACATTAAAGTTTCCAGAAATAACACAGTAAAGAAA  
TATGAAAATAAACTGGAAAATAAAATATACCCACCCATCCGAAAAATCTACATCATCTCT  
TTCATTTGTCCCAATGCCTTTC

Quality: High quality sequence stops at base: 344

Entry Created: Oct 15 1997  
Last Updated: Mar 2 1998

## COMMENTS

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further information.

## LIBRARY

Lib Name: Soares\_testis\_NHT  
Organism: Homo sapiens  
Sex: male  
Lab host: DH10B  
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker  
R. Site 1: Not I  
R. Site 2: Eco RI  
Description: 1st strand cDNA was prepared from mRNA obtained from Clontech Laboratories, Inc., and primed with a Not I - oligo(dT) primer [5' TGTTACCAATCTGAAGTGGGAGCGGCCGCCCAATTTTTTTTTTTTTTTTTT 3']. Double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization to Cot5, and was constructed by Bento Soares and M. Fatima Bonaldo.

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

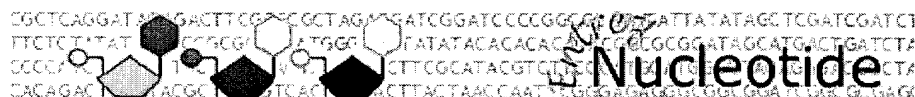
Title: WashU-NCI human EST Project  
Authors: Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R., Wilson,R.  
Year: 1997  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



Boo

Go

Clear

## Details

Show:

**Send to**

File 

## Links

```

Lib Name:      NCI_CGAP_Ut3
Organism:      Homo sapiens
Organ:         uterus
Tissue type:   poorly-differentiated endometrial adenocarcinoma, 2 pooled
                tumors
Lab host:      DH10B

```

Vector: pCMV-SPORT6  
R. Site 1: SalI  
R. Site 2: NotI  
Description: Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.45 kb. Life Technologies catalog #: 11541-018

**SUBMITTER**

Name: Robert Strausberg, Ph.D.  
E-mail: [cgapbs-r@mail.nih.gov](mailto:cgapbs-r@mail.nih.gov)

**CITATIONS**

Title: National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index  
Authors: NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>  
Year: 1997  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



**Nucleotide**

PubMed

Nucleotide

Protein

Genome

## Structure

PMC

## Taxonomy

OMIM

Boo

## Search

Nucleotide

for

for \_\_\_\_\_

Go

Clear

## Limits

[Preview/Index](#)

## History

Clipboard

## Details

## Display

default

Show:

20 

Send to

File

□ 1: N52026. yz08e07.s1 Soares...[gi:1193192]

## Links

## IDENTIFIERS

```
dbEST Id: 457555
EST name: yz08e07.s1
GenBank Acc: N52026
GenBank gi: 1193192
GDB Id: 3900996
```

## CLONE INFO

Clone Id: IMAGE:282468 (3')  
Source: IMAGE Consortium, LLNL  
DNA type: cDNA

## PRIMERS

Sequencing: m13 -40 forward  
PolyA Tail: Unknown

## SEQUENCE

ACCCTGAGATAGAATAGATTTATTAGCAAGAGGTAATCAGGAAACATATATTTTTTACAA  
AATGGAAATTTTTTTTCCAAACAAGCTGTAAGTTGAAATATTTTAGGACTTGAAATAGAAT  
TCTCATACCACTAGGTATTGCTTACAGCAAAAGTTGTCTGTCTGTTGTAGTGGAGCATGC  
CTGCCACTTCGGAGTTAACCTGTGTTTTCTATACTGTACAGTGTAaaaaATACATGGTAA  
TATTCACAGAATAAGCACTACATTACTATATTCTGTCTAGAAGGCATTTAGACAGGACTA  
CAGTATATGCCATAAAAACACTTGGTTATTGGATTTTCCCTAATTCTACAGTGTGGGTA  
CTAAATTATCCACCAAGGTATACNGGACTTAAGAGCCATCCTCAATGGTAAGGCCTGGTA  
AGTGACCCGTTANGCAGCCTTAAAGANGGAAAAAGTGACATTTTGGGGTTCCCCGACTT  
TCAGTG

Quality: High quality sequence stops at base: 194

Entry Created: Feb 15 1996

Last Updated: Feb 15 1996

## COMMENTS

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further information.

## LIBRARY

```

Lib Name:      Soares_multiple_sclerosis_2NbHMSP
Organism:      Homo sapiens
Sex:           male
Tissue type:   multiple sclerosis lesions
Develop. stage: Age 46
Lab host:      DH10B (ampicillin resistant)
Vector:        pT7T3D (Pharmacia) with a modified polylinker V_TYPE:
                phagemid
R. Site 1:     Not I
R. Site 2:     Eco RI

```

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer [5' TGTTACCAATCTGAAGTGGGAGCGGCCGCATTTTTTTTTTTTTTTTTTTT 3'], double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from 4 multiple sclerosis lesions from one patient was kindly provided by Dr. Kevin G. Becker (NINDS/NIH).

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

Title: The WashU-Merck EST Project  
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfig,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P., Wilson,R.  
Year: 1995  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45





CGCTCAGGATATGACTTCCTCTCCTAGATCGGATCCCGGGGATTATTATATAGCTCGATCGATC  
TTCTCTATATCCGGGATGCTGCTATATACACACACACCGCGCGGATAGCATCACTGATCT  
CCCCATCT  
CACAGACCTACGCT

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

OMIM

Boo

Search

Nucleotide

for

Go

Clear

Limits

Preview/Index

History

Clipboard

Details

Display

default

Show:

20

Send to

File

☐ 1: H98215.yx09a02.s1 Soares...[gi:1119100]

Links

## IDENTIFIERS

dbEST Id: 407094  
EST name: yx09a02.s1  
GenBank Acc: H98215  
GenBank gi: 1119100  
GDB Id: 3870836

## CLONE INFO

Clone Id: IMAGE:261194 (3')  
Insert length: 1053  
DNA type: cDNA

## PRIMERS

Sequencing: m13 -40 forward  
PolyA Tail: Unknown

## SEQUENCE

CACAGGAACAATTCTTTTATTGTACATTGGAGAAATAGCCCTGTGTGCTGGTTCAAGGTG  
CAACATACAGAATATTGAATTAAGAAAAGAGGGAACGGGGAAGGGAANGGAAACCTCTTT  
GAGGTCCAAAGTTGNCAACAAAAAATGGTAAAAGATTTCTCACGCAAGANGGCATTTTT  
GCAATACCATGCAAAACAGGCAGCTGGTGTGCCTTAAGAGAATCCCTATAAATAACAGA  
AAAGACACTCCAAGCATTCTGTACGTGGACTCAGAGCACAGAGAAAAGAACTAAAATG  
CCTTTTGGATTTCAAGATATTTGGCACTCTTGTGATTACATTTTTTTTACAGTCCATTAAA  
GGGAATAAACTGACATAATATT

Quality: High quality sequence stops at base: 109

Entry Created: Dec 12 1995  
Last Updated: Dec 12 1995

## COMMENTS

High quality sequence stops: 109  
Source: IMAGE Consortium, LLNL  
This clone is available royalty-free through LLNL ; contact  
the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for further  
information.

## LIBRARY

Lib Name: Soares melanocyte 2NbHM  
Organism: Homo sapiens  
Sex: Male  
Tissue type: melanocyte  
Lab host: DH10B (ampicillin resistant)  
Vector: pT7T3D (Pharmacia) with a modified polylinker  
R. Site 1: Not I  
R. Site 2: Eco RI  
Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer  
[5' TGTTACCAATCTGAAGTGGGAGCGGCCGAGTTTTTTTTTTTTTTTTTTT 3'],

double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from normal foreskin melanocytes (FS374) was kindly provided by Dr. Anthony P. Albino.

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

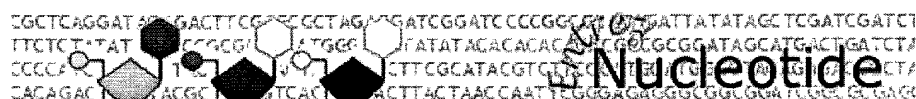
Title: The WashU-Merck EST Project  
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,  
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,  
Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,  
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,  
Wilson,R.  
Year: 1995  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



Bon

Go Clear

## Details

**Send to**

## Links

dbEST Id: 157681  
EST name: yd83b10.r1  
GenBank Acc: T87560  
GenBank gi: 715912  
GDB Id: 470428

Clone Id: IMAGE:114811 (5')  
Insert length: 1110  
DNA type: cDNA

Sequencing: M13RP1  
PolyA Tail: Unknown

NTAATACTGAGAGCGGAAGATTTTTTTAAAGAGATATGGTAGTTTAAATAGGTTTGTTTA  
AAGAACTCTTGGGAAATTTCTTAACAAATTGTAATTGTTTTTATAAACCTTCCTACAGAG  
TTGTGTACCATTTCATCATCACCATCATTTAATATCTGTAAGTGCCAGAAAGGTACAAGGT  
GCTATACAGCCGTTAAACACCCCTCCCTGCAACTGTTTAAAATTAGAATAGATTTTTTGC  
ATCAACCACTGCTAGAAGCTTTCCCCACCTAGGTGGCCTGGCAAGGGCCTGCATTCCCCAA  
CATCCCATTCATATGTTTCCCACTCCAGAAAGTCTCTGGGGTTTATTCAGCANTTTTTATC  
NTGGACGTGTCCCNTTTACCCCTGGGTTTGGGTTGGGTTGGGTTTGTTTTTTAAANTC  
AGGAAAGTTTTTCAGGGATTCAAAGGGGGTCNGACNTTTTTTAAACCGGCGGAGTACAGG  
TCNTTTTTTTTGCTANGGTTTTGTGTGGTACAGNAACCCANTTGGCCNTTAGTGTGT  
GGGTATTTTGT

Quality: High quality sequence stops at base: 324

Entry Created: Mar 17 1995  
Last Updated: Mar 17 1995

Insert Size: 11110  
High quality sequence stops: 324 Source: IMAGE Consortium,  
LLNL This clone is available royalty-free through LLNL ;  
contact the IMAGE Consortium ([info@image.llnl.gov](mailto:info@image.llnl.gov)) for  
further information.

Lib Name: Soares fetal liver spleen 1NFLS  
Organism: Homo sapiens  
Sex: male  
Organ: Liver and Spleen  
Develop. stage: 20 week-post conception fetus  
Lab host: DH10B (ampicillin resistant)  
Vector: pT7T3D (Pharmacia) with a modified polylinker

R. Site 1: Pac I  
R. Site 2: Eco RI  
Description: 1st strand cDNA was primed with a Pac I - oligo(dT) primer  
[5' AACTGGAAGAATTAATTAAAGATCTTTTTTTTTTTTTTTTTTTT 3'],  
double-stranded cDNA was ligated to Eco RI adaptors  
(Pharmacia), digested with Pac I and cloned into the Pac I  
and Eco RI sites of the modified pT7T3 vector. Library went  
through one round of normalization. Library constructed by  
Bento Soares and M.Fatima Bonaldo.

**SUBMITTER**

Name: Wilson RK  
Institution: Washington University School of Medicine  
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108  
Tel: 314 286 1800  
Fax: 314 286 1810  
E-mail: [est@watson.wustl.edu](mailto:est@watson.wustl.edu)

**CITATIONS**

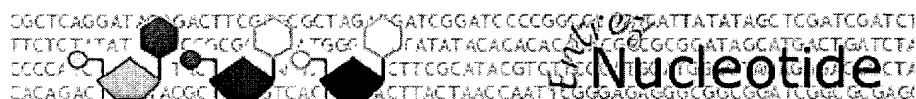
Title: The WashU-Merck EST Project  
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,  
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.,  
Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,  
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,  
Wilson,R.  
Year: 1995  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



Boo

Go

## Clear

## Details

## Display

default

Show:

20

**Send to**

File

## Links

dbEST Id: 3658251  
EST name: xv76c12.x1  
GenBank Acc: AW275150  
GenBank qi: 6662180

Clone Id: IMAGE:2819062 (3')  
Source: NCI  
DNA type: cDNA

Sequencing: -40UP from Gibco  
PolyA Tail: Unknown

TTTTTTTGGAGGAATGAGCAATTTATTTCAATAAAGAGAAAGCATTAAATTTTGCTACAGTG  
GGAAAAAATGAACTCAAGAGTTGCTACATTTAACTGTATCCCCATTTATCTCTGCACGAT  
GTCTTTATCTCAGTGTCTCAATTCACACTAAAATATTGAATGAGAAATACACCACGTTGG  
CTGATTGCTTGACATGTCTGATTTAGGGGAGACTTCTACAACCACTCCTCTCTTTTTTCTC  
CCAGTAAATACTTTTGACTTTGACACCTACCATATTGGAAATGACAGGTGCCCGAGGGCA  
AGTGCATCAAAGCAGTTAGGATTCTGAATGCTTGCTAAGGATTATTTTTTTAATGGAGCAG  
TTCTATTGAATCCTGAAAAATGCAGAACCCAGTGAATCTCCTTTAT

Quality: High quality sequence stops at base: 404

Entry Created: Jan 3 2000

Last Updated: Jan 3 2000

Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The I.M.A.G.E. Consortium DNA Sequencing by: Washington University Genome Sequencing Center  
Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: [www-bio.llnl.gov/bbrp/image/image.html](http://www-bio.llnl.gov/bbrp/image/image.html)

```

Lib Name:      NCI_CGAP_Lu28
Organism:      Homo sapiens
Organ:         lung
Tissue type:   two pooled squamous cell carcinomas
Lab host:      DH10B
Vector:        pCMV-SPORT6
R. Site 1:     SalI
R. Site 2:     NotI

```

Description: Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies.

**SUBMITTER**

Name: Robert Strausberg, Ph.D.  
E-mail: [cgapbs-r@mail.nih.gov](mailto:cgapbs-r@mail.nih.gov)

**CITATIONS**

Title: National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index  
Authors: NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>  
Year: 1997  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45



CGCTCAGGATATGACTTCGTCGCTAGAGATCGGATCCCGGGGATGATTATATAGCTCGATCGATC  
ITCTCTATATTCGGGATATGGGATATATACACACACATCGCGCGGATAGCATGACTGATCTA  
CCCCATCTATTCGCTATCTCTCGCATACGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT  
CACAGACTCTACGCT

PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

OMIM

Bio

Search **Nucleotide** for

Go

Clear

Limits

Preview/Index

History

Clipboard

Details

Display

default

Show:

20

Send to

File

☐ 1: AI521564. to64g08.x1 NCI\_CG...[gi:4435699]

Links

**IDENTIFIERS**

dbEST Id: 2337724  
EST name: to64g08.x1  
GenBank Acc: AI521564  
GenBank gi: 4435699

**CLONE INFO**

Clone Id: IMAGE:2183102 (3')  
Source: NCI  
Insert length: 4578  
DNA type: cDNA

**PRIMERS**

Sequencing: -40UP from Gibco  
PolyA Tail: no

**SEQUENCE**

AGCGGCCGCCCTTTTTTTTTTTTTTTTTTTTTTTTGGAGGAATGAGCAATTTATTTCAATAAAG  
AGAAAGCATTAATTTTGCTACAGTGGGAAAAAATGAACTCAAGAGTTGCTACATTTAACT  
GTATCCCCATTTATCTCTGCACGATGTCTTTATCTCAGTGTCTCAATTCACACTAAAATA  
TTGAATGAGAAATACACCACGTTGGCTGATTGCTTGACATGTCTGATTTAGGGGAGACTTC  
TACAACCACTCCTCTCTTTTTTCTCCAGTAAATACTTTTGACTTTGACACCTACCATAT  
TGGAAATGACAGGTGCCCCAGGGCAAGTGCATCAAAGCAGTTAGGATTCTGAATGCTTGCT  
AAGGATTATTTTTTAAATGGAGCAGTTCTATTGAATCCTGAAAATGCAGAACCCAG

Quality: High quality sequence stops at base: 371

Entry Created: Mar 16 1999  
Last Updated: May 12 1999

**COMMENTS**

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,  
Michael R. Emmert-Buck, M.D., Ph.D.  
cDNA Library Preparation: Life Technologies, Inc.  
cDNA Library Arrayed by: Greg Lennon, Ph.D.  
DNA Sequencing by: Washington University Genome Sequencing  
Center  
Clone distribution: NCI-CGAP clone distribution information  
can be found through the I.M.A.G.E. Consortium/LLNL at:  
[www-bio.llnl.gov/bbrp/image/image.html](http://www-bio.llnl.gov/bbrp/image/image.html)

**LIBRARY**

Lib Name: NCI\_CGAP\_Gas4  
Organism: Homo sapiens  
Organ: stomach  
Tissue type: poorly differentiated adenocarcinoma with signet ring cell  
features  
Lab host: DH10B

Vector: pCMV-SPORT6  
R. Site 1: SalI  
R. Site 2: NotI  
Description: Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.69 kb. Life Technologies catalog #: 11549-011

**SUBMITTER**

Name: Robert Strausberg, Ph.D.  
E-mail: [cgapbs-r@mail.nih.gov](mailto:cgapbs-r@mail.nih.gov)

**CITATIONS**

Title: National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index  
Authors: NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>  
Year: 1997  
Status: Unpublished

**MAP DATA**

---

[Disclaimer](#) | [Write to the Help Desk](#)  
[NCBI](#) | [NLM](#) | [NIH](#)

Jun 19 2003 12:37:45